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Inventor(s) Cannon et al.
For: "Human Intestinal Npt2B"
Docket No. ROCH-001DIV

FIG. 1

1 MAPWPELGDA QPNPDKYLEG AAGQQPTAPD KSKETNKNNT EAPVTKIELL
51 PSYSTATLID EPTEVDDPWN LPTLQDSGIK WSERDTKGKI LCFFQGIGRL
101 ILLLGFYFF VCSLDILSSA FQLVGGKMAG QFFSNSSIMS NPLLGLVIGV
151 LVTVLVQSSS TSTSIVVSMV SSSLTVRAA IPIIMGANIG TSITNTIVAL
201 MQVGDRSEFR RAFAGATVHD FFNWLSVLVL LPVEVATHYL EIITQLIVES
251 FHFKNGEDAP DLLKVITKPF TKLIVQLDKK VISQIAMNDE KAKNKS LVKI
301 WCKTFTNKTQ INVTPSTAN CTSPSLCWTD GIQNWTMKNV TYKENIAKCQ
351 HIFVNFHLPD LAVGTILLIL SLLVLCGCLI MIVKILGSVL KGQVATVIKK
401 TINTDFPFPF AWLTGYLAIL VGAGMTFIVQ SSSVFTSALT PLIGIGVITI
451 ERAYPLTLGS NIGTTTTAIL AALASPGNAL RSSLQIALCH FFFNISGILL
501 WYPIPFTRLP IRMAKGLGNI SAKYRWFAVF YLIIFFFLIP LTVFGLSLAG
551 WRVLVGVGVP VVFIIILVLC LRLLQSRCPR VLPKKLQNWN FLPLWMRSLK
601 PWDAVVSKFT GCFQMRCCCC CRVCCRACCL LCGCPKCCRC SKCSEDLEEA
651 QEGQDVVK PETFDNITIS REAQGEVPAS DSKTECTAL* (SEQ ID NO:01)

FIG. 2i

1 CTGACGTAGG CCCAGCACCT GCGGAGGGAG CGCTGACCAT GGCTCCCTGG
51 CCTGAATTGG GAGATGCCA GCCCAACCCC GATAAGTACC TCGAAGGGC
101 CGCAGGTCAAG CAGCCCCTG CCCCTGATAA AAGCAAAGAG ACCAACAAAA
151 ATAACACTGA GGCACCTGTA ACCAAGATTG AACTTCTGCC GTCTACTCC
201 ACGGCTACAC TGATAGATGA GCCCACTGAG GTGGATGACC CCTGGAACCT
251 ACCCACTCTT CAGGACTCGG GGATCAAGTG GTCAGAGAGA GACACCAAAG
301 GGAAGATTCT CTGTTTCTTC CAAGGGATTG GGAGATTGAT TTTACTCTC
351 GGATTTCTCT ACTTTTCGT GTGCTCCCTG GATATTCTTA GTAGCGCCTT
401 CCAGCTGGTT GGAGGAAAAA TGGCAGGACA GTTCTTCAGC AACAGCTCA
451 TTATGTCCAA CCCTTTGTTG GGGCTGGTGA TCGGGGTGCT GGTGACCGTC
501 TTGGTGCAGA GCTCCAGCAC CTCAACGTCC ATCGTTGTCA GCATGGTGT
551 CTCTTCATTG CTCACTGTTC GGGCTGCCAT CCCATTATC ATGGGGCCA
601 ACATTGGAAC GTCAATCACC AACACTATTG TTGCGCTCAT GCAGGTGGG
651 GATCGGAGTG AGTTCAGAAG AGCTTTGCA GGAGCCACTG TCCATGACTT
701 CTTCAACTGG CTGTCCTGT TGTTGCTCTT GCCCCTGGAG GTGGCCACCC
751 ATTACCTCGA GATCATAACC CAGCTTATAG TGGAGAGCTT CCACTTCAAG
801 AATGGAGAAG ATGCCCGAGA TCTTCTGAAA GTCATCACTA AGCCCTTCAC
851 AAAGCTCATT GTCCAGCTGG ATAAAAAAAGT TATCAGCCAA ATTGCAATGA
901 ACGATGAAAA AGCGAAAAAC AAGAGTCTTG TCAAGATTG GTGCAAAACT
951 TTTACCAACA AGACCCAGAT TAACGTCACT GTTCCCTCGA CTGCTAAGT
1001 CACCTCCCCCT TCCCTCTGTT GGACGGATGG CATCCAAAAC TGGACCATGA
1051 AGAATGTGAC CTACAAGGAG AACATGCCA AATGCCAGCA TATCTTGTG
1101 AATTTCACC TCCCCTGGATCT TGCTGTGGC ACCATCTTGC TCATACTCTC
1151 CCTGCTGGTC CTCTGTGGTT GCCTGATCAT GATTGTCAAG ATCCTGGCT
1201 CTGTGCTCAA GGGGCAGGTC GCCACTGTCA TCAAGAAGAC CATCAACACT
1251 GATTTCCCCCT TTCCCTTTGC ATGGTTGACT GGCTACCTGG CCATCCTCGT
1301 CGGGGCAGGC ATGACCTTCA TCGTACAGAG CAGCTCTGTG TTCACGTCGG
1351 CCTTGACCCC CCTGATTGGA ATCGGGGTGA TAACCATTGA GAGGGCTTAT
1401 CCACTCACGC TGGGCTCCAA CATCGGCACC ACCACCACCG CCATCCTGGC
1451 CGCCTTAGCC AGCCCTGGCA ATGCATTGAG GAGTCACTC CAGATGCC
1501 TGTGCCACTT TTTCTTCAAC ATCTCCGGCA TCTTGCTGTG GTACCCGATC
1551 CCGTTCACTC GCCTGCCCAT CCGCATGGCC AAGGGCTGG GCAACATCTC
1601 TGCCAAGTAT CGCTGGTTCG CCGTCTTCTA CCTGATCATC TTCTTCTTCC
1651 TGATCCCGCT GACGGTGTG TGCTCTCGC TGGCCGGCTG GCGGGTGTG
1701 GTTGGTGTG TGTTCCCGT CGTCTTCATC ATCATCCTGG TACTGTGCCT
1751 CCGACTCCTG CAGTCTCGCT GCCCACCGT CCTGCCGAAG AAAACTCCAGA
1801 ACTGGAACTT CCTGCCGCTG TGGATGCGCT CGCTGAAGCC CTGGGATGCC
1851 GTCGTCCTCA AGTTCACCGG CTGCTTCCAG ATGCGCTGCT GCTGCTGCTG
1901 CGCGCTGTG TGCCGCGCGT GCTGTTGCT GTTGGCTGC CCCAAGTGC
1951 CGCGCTGCAG CAAGTGTGCA GAGGACTTGG AGGAGGGCGCA GGAGGGCAG
2001 GATGTCCCTG TCAAGGCTCC TGAGACCTT GATAACATAA CCATTAGCAG
2051 AGAGGCTCAG GGTGAGGTCC CTGCCTCGGA CTCAAAGACC GAATGCACGG
2101 CCTTGTAGGG GACGCCAGG ATTGTCAAGG ATGGGGGGAT GGTCTTGAG
2151 TTTTGCATGC TCTCTCCCT CCCACTTCTG CACCTTTCA CCACCTCGAG
2201 GAGATTGCT CCCATTAGC GAATGAAATT GATGCAGTCC TACCTAACTC
2251 GATTCCTTT GGCTTGGTGG GTAGGCCCTGC AGGGCACTTT TATTCCAACC
2301 CCTGGTCACT CAGTAATCTT TTACTCCAGG AAGGCACAGG ATGGTACCTA
2351 AAGAGAATTA GAGAATGAAC CTGGCGGGAC GGATGTCTAA TCCTGCACCT
2401 AGCTGGGTTG GTCAAGTAGAA CCTATTTCA GACTAAAAA CCATCTTCAG
2451 AAAGAAAAGG CCCAGGGAAAG GAATGTATGA GAGGCTCTCC CAGATGAGGA
2501 AGTGTACTCT CTATGACTAT CAAGCTCAGG CCTCTCCCTT TTTTAAACC
2551 AAAGTCTGGC AACCAAGAGC AGCAGCTCCA TGGCCTCCTT GCCCCAGATC
2601 AGCCTGGTC AGGGGACATA GTGTCAATTGT TTGGAAACTG CAGACCACAA

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FIG. 2ii

2651 GGTGTGGGTC TATCCCACCTT CCTAGTGCTC CCCACATTCC CCATCAGGGC
2701 TTCCCTCACGT GGACAGGTGT GCTAGTCCAG GCAGTTCACT TGCAGTTCC
2751 TTGTCCTCAT GCTTCGGGGA TGGGAGGCCAC GCCTGAACTA GAGTCAGGC
2801 TGGATACATG TGCTCACCTG CTGCTCTTGT CTTCTAAGA GACAGAGAGT
2851 GGGGCAGATG GAGGAGAAGA AAGTGAGGAA TGAGTAGCAT AGCATTCTGC
2901 CAAAAGGGCC CCAGATTCTT AATTAGCAA ACTAAGAAC CCAATTCAAA
2951 AGCATTGTGG CTAAAGTCTA ACGCTCCTCT CTTGGTCAGA TAACAAAAAGC
3001 CCTCCCTGTT GGATCTTTG AAATAAAACG TGCAAGTTAT CCAGGCTCGT
3051 AGCCTGCATG CTGCCACCTT GAATCCCAGG GAGTATCTGC ACCTGGAATA
3101 GCTCTCCACC CCTCTCTGCC TCCTTACTTT CTGTGCAAGA TGACTTCCTG
3151 GTTAACTTC CTTCTTTCCA TCCACCCACC CACTGGAATC TCTTCCAAA
3201 CATTTTCCA TTTCCCACA GATGGGCTTT GATTAGCTGT CCTCTCTCCA
3251 TGCCTGCAAA GCTCCAGATT TTTGGGGAAA GCTGTACCCA ACTGGACTGC
3301 CCAGTGAACG GGGATCATTG AGTACAGTCG AGCACACGTG TGTGCATGGG
3351 TCAAAGGGGT GTGTTCCCTC TCATCCTAGA TGCCCTCTCT GTGCCCTCCA
3401 CAGCCTCCTG CCTGATTACA CCACTGCCCG CGCCCCACCC TCAGCCATCC
3451 CAATTCTTCC TGGCCAGTGC GCTCAGCCT TATCTAGGAA AGGAGGAGTG
3501 GGTGTAGCCG TGCAGCAAGA TTGGGGCTC CCCCATCCCA GCTTCTCCAC
3551 CATCCCAGCA AGTCAGGATA TCAGACAGTC CTCCTCTGAC CCTCCCCCTT
3601 GTAGATATCA ATTCCCAAAC AGAGCCAAAT ACTCTATATC TATAGTCACA
3651 GCCCTGTACA GCATTTTCA TAAGTTATAT AGTAAATGGT CTTCTAGTGC
3701 TCTCATTTGG AAATGAGGCA GGCTTCTTCT ATGAAATGTA AAGAAAGAAA
3751 CCACTTGTA TATTTGTA TACCACCTCT GTGGCCATGC CTGCCCGGCC
3801 CACTCTGTAT ATATGTAAGT TAAACCCGGG CAGGGGCTGT GGCGTCTTT
3851 GTACTCTGGT GATTTTTAGA AATTGAATCT TTGTAATTGC ATTGATTGTA
3901 TAATAATTTC GAGACCAGGT CTCGCTGTGT TGCTCAGGCT GGTCTCAAAC
3951 TCCTGAGATC AAGCAATCCG CCCACCTCAG CCTCCCAAAG TGCTGAGATC
4001 ACAGGGGTGA GCCACCACCA GGCCTGATTG TAATTTTTT TTTTTTTTT
4051 TTTACTGGTT ATGGGAAGGG AGAAATAAAA TCATCAAACC CAAAAAA
4101 AAAAAAAAAA AAAAAAAAAA AAAAAAAAAA AAAAAAA (SEQ ID NO:02)